

- 1 -

piece 1, NC_000913, accD_dedA+, config: linear, direction: +, begin: 2431919, end: 2432123

Sequence alignment of nine homologous proteins. The proteins are aligned along a horizontal axis, with their sequence labels above them. A red dot highlights a conserved cysteine residue at position 1990 in all proteins except the first one. The first protein has a valine at this position. The second protein has a leucine. The third protein has a leucine. The fourth protein has an arginine. The fifth protein has an aspartic acid. The sixth protein has a serine. The seventh protein has a valine. The eighth protein has a leucine. The ninth protein has a threonine.

1 NC 000813 accD

μ₀

35 3.0 bits

{--- ... sd-(8)=ir 2432006 Gap

• [View all \(1\)](#) [Recent posts](#)

|--- ... sd-dir 2432006 accD_dedA+
|--- ... p35-(24)-p10 2432003 Gap
|--- ... p35-p10 2432003 total 6 0

```
sd-8) ir 2432006 Gap 2.4 bits  
sd-ir 2432006 accD_dedA+ total 7.9 bits  
p35-(24)-p10 2432003 Gap 2.4 bits  
p35-p10 2432003 total 6.0 bits  
ir accD_dedA+
```

25 / 241 10.2.102010

1 bits

- 15

புதுப்பு p10 5.5 bits

* * * * * 24232090 * * * 24232100 * * * 24232110 * * * 24232120

5' t g g t c g a a c c g c g a g t t a c t t t t t a t t t g c g c g c g c g c g 3'

-trp - ser - asn - arg - gly - val - thr - phe - tyr - phe - ala - ala - arg - ala -
- - - - - - - - - - - - - -fMet - arg - arg - val - pro -
- - - val - glu - pro - arg - ser - tyr - phe - leu - phe - cys - gly - ala - cys -

p35 2.5 bits

 p10 5.2 bits

{-----} p35-(24)-p10 2432106 Gap 2.4 bits
-----| p35-p10 2432106 total 5.3 bits